\_\_\_\_\_\_

Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Fri Aug 31 11:15:03 EDT 2007

\_\_\_\_\_\_

\*\*\*\*\*\*\*\*\*\*\*\*\*

Reviewer Comments:

<210> 55

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<400> 55

Asp Leu Gln Leu Arg Leu

1 5

The above <223> response is incorrect: this is not a DNA sequence.

## Validated By CRFValidator v 1.0.3

Application No: Version No: 10574470 1.0

Input Set:

Output Set:

**Started:** 2007-08-31 10:46:26.824 Finished:

2007-08-31 10:46:29.736

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 912 ms

Total Warnings: 152

Total Errors:

No. of SeqIDs Defined: 164

> Actual SeqID Count: 164

Error code		Error Description									
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W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(2)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(3)
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W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(17)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(20)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(21)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(22)

Input Set:

Output Set:

**Started:** 2007-08-31 10:46:26.824

Finished: 2007-08-31 10:46:29.736

**Elapsed:** 0 hr(s) 0 min(s) 2 sec(s) 912 ms

Total Warnings: 152
Total Errors: 0

No. of SeqIDs Defined: 164

Actual SeqID Count: 164

**Error Description** 

Error code

This error has occured more than 20 times, will not be displayed

## SEQUENCE LISTING

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     National Institute of Advanced Industrial Science and Technology
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<151> 2004-01-07
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Leu Asp Leu Asn Leu Ala Pro Pro Met Glu Phe

10

5

1

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20 25 30

Asp His Asp Tyr Leu Leu Gly Phe Ser Trp Pro Pro Arg Ser Tyr Thr 35 40 45

Cys Ser Phe Cys Lys Arg Glu Phe Arg Ser Ala Gln Ala Leu Gly Gly 50 55 60

His Met Asn Val His Arg Arg Asp Arg Ala Arg Leu Arg Leu Gln Gln 65 70 75 80

Ser Pro Ser Ser Ser Thr Pro Ser Pro Pro Tyr Pro Asn Pro Asn 85 90 95

Tyr Ser Tyr Ser Thr Met Ala Asn Ser Pro Pro Pro His His Ser Pro
100 105 110

Leu Thr Leu Phe Pro Thr Leu Ser Pro Pro Ser Ser Pro Arg Tyr Arg
115 120 125

Ala Gly Leu Ile Arg Ser Leu Ser Pro Lys Ser Lys His Thr Pro Glu 130 135 140

Ala Thr Arg Phe Thr Ser Lys Asp Ala Cys Lys Ile Leu Arg Asn Asp 165 170 175

Glu Ile Ile Ser Leu Glu Leu Glu Ile Gly Leu Ile Asn Glu Ser Glu 180 185 190

Gln Asp Leu Asp Leu Glu Leu Arg Leu Gly Phe Ala 195 200

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Ser Glu Gln Asp Leu Asp Leu Glu Leu Arg Leu Gly Phe Ala
20 25 30

<210> 20

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<212> PRT

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1 5
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<210> 34

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 1
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 1
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